# UMass CS690N: Advanced Natural Language Processing, Spring 2017

## Assignment 1

Due: Feb 10

**Introduction:** This homework includes a few mathematical derivations and an implementation of one simple language model—Saul and Pereira's "aggregate" latent-class bigram model. Pereira (2000) uses the model to try to disprove one of the most famous foundational examples in 20th century linguistics; you will replicate this experiment and assess the evidence. (Pereira 2000, "Formal grammar and information theory: together again?," Phil. Trans. R. Soc. Lond. A, 358, 1239-1253.)

You can implement code with whatever language you like; we recommend Python, but other math-friendly dynamic languages are fine (Matlab, Julia, maybe R though it's slow); or really anything like Scala, Java, Lua, Haskell, C++ or whatever.

- Write your final answers in a document and upload as a PDF (to Gradescope).
- Upload a zip file of your code (to Moodle). Do NOT include data files.
- For the writeup:
  - Put each full question on a new page (necessary for Gradescope!)
  - Clearly mark the numbers of each question you're answering.
  - Report natural logarithms for all questions.

**Data:** Please use the Brown corpus from the NLTK download at http://www.nltk.org/nltk\_data/. Strip out POS tags and lowercase all words. There should be 57340 sentences, 1161169 tokens (not including START or END symbols) and vocabulary size 49740 (not including END, though it may be convenient to throw in START and END into the vocabulary). Note the END symbols are NOT the same thing as sentence-final punctuation; punctuation symbols are themselves tokens. (Not all sentences end with punctuation!) The Brown Corpus defines tokenization and sentence boundaries which we'll be using.

Take a look at some of the files so you have a sense of what's in it. The Brown Corpus was a pioneering dataset when it was released in the 1960s. Though it's tiny by modern standards (1M words), it's big enough to do a little latent-variable language learning.

To help verify it's loaded correctly, try printing out for yourself the most and least common words; for example, with python's collections. Counter and its most\_common() (or just use a dict then sorted(dict.items(), key=lambda (w,c): -c)[:10]). You will use these little counting/ranking tricks all the time in NLP.

### Q1: Models

**Q1.1** (Unigram model; 2 points): Calculate and report  $\log p$  ("colorless green ideas sleep furiously END") under an MLE unigram model.

Technically speaking, when evaluating a sentence's probability, you should always include the probability of generating the END token; this yields a proper, sums-to-1 probability distribution over all possible sentences.

**Q1.2** (**Bigram model**; **1 point**): What is the probability of this sentence under an MLE bigram model (a.k.a. first order Markov model)? Why?

Q1.3 (8 points): List the parameters for Saul and Pereira's "aggregate bigram" model. Give them variable names and say, in short English descriptions, what the variables mean. How many parameters total does the full model have (the total dimensionality)? Notes: 1) the z variables are not considered parameters for this purpose, and 2) You can have fun thinking about whether to count the last dimension in a multinomial parameter vector, because technically, it's extraneous. We don't care whether you include it or not for this problem.

**Q1.4** (2 points): Assume the model is learned and you have access to the parameters. In terms of the parameters you have defined, write out the equation that can be calculated to evaluate the log-probability of a new sentence  $(w_1, \ldots, w_T)$ :

$$\log p(w_1, w_2, \dots, w_{T+1} = \text{END} \mid w_0 = \text{START})$$

Technical note: In the above equation, we're trying to be careful with START/END symbols. Tokens 1 through T are actual observed word tokens; the word at {T+1} isn't really a word, but a dummy symbol that indicates the sentence isn't ending. START also isn't a word, and the model also doesn't generate it, but assumes it's already there.

### **Q2: END.**

**Q2.1** (5 points): Why is it necessary to model the END symbol? For example, in order to use a language model for translation or something, you could instead calculate the following equation:

$$p(w_1, w_2, \dots, w_T \mid w_0 = \text{START})$$

Please give a reason why modeling END is important for linguistic quality. Use an example to back up your argument.

**Q2.2** (5 points): Consider a different reason to model END: it is necessary for a well-formed probability distribution over all strings. (Formally, a string is a sequence of any integer length of symbols from a fixed set of symbols (a.k.a. vocabulary)  $\mathcal{V}$ .) A well-formed probability distribution is one that obeys the usual axioms of probability. Show that if you use the equation in Q2.1 to calculate a string's probability, the probability distribution you define is not well-formed.

• Hint: "show" means "prove" (that is, argue mathematically). A correct answer can be very short.

### Q3: EM.

**Q3.1** (5 points): Write the E-step update equation for Saul and Pereira's model, in pseudocode that is mathematically precise, uses the notation you've defined, but is easier to read than computer code. You will have to define new mathematical variables; explain in English what they mean.

**Q3.2** (5 points): Write the M-step update equations for the model in a similar manner.

## Q4: Implementing EM.

Implement Saul and Pereira's latent-class transition model with EM training on the Brown corpus. EM can be tricky to debug; a set of implementation tips are included at the end of this document. Read over the questions below before starting; they ask about pieces of your implementation which you may be able to implement incrementally, instead of all at once.

**Algorithm:** We suggest using the following order for the algorithm. You can combine the E-step and the M-step 1 if desired (because you don't need to store the token-level posteriors).

- Randomly initialize the parameters.
- For each iteration:
  - (E step): Calculate posteriors at each token level  $p(z_t \mid w_{t-1}, w_t)$ .
  - (M-step phase 1): Calculate expected counts of transitions and emissions.
  - (M-step phase 2): Normalize these counts so they correctly sum to 1 to be the new parameters (note normalization is different for each parameter matrix).

**Verification of M-step counts:** You have to calculate summary statistics at the start of the M-step: the expected counts of word-to-state transitions (a matrix  $V \times K$ ), and the expected counts of the state-to-word emissions (a matrix  $V \times K$ ). One way to debug the E-step is to confirm that, if you go by the matrix shapes above, the row sums equal the raw word counts in the data. (You of course don't need to implement them with matrixes exactly like this; this is for illustration.)

**Q4.1** (1 point): Explain why this should be the case.

**Q4.2** (4 points): Check whether this is the case empirically in your code. If the numbers are way different, there may be a bug in your code. Once you're reasonably certain you're debugged, report a little bit whether the numbers are the same or some of the numbers are the same. Give a useful quantitative and/or qualitative summary to compare these vectors.

**Q4.3** (Verification of M-step probabilities; 2 points): Verify that your parameters sum to 1 the way you think they should. Explain.

**Q4.4** (Verification of EM as an optimizer; 8 points): It can be proved that, as EM is running, the marginal log-likelihood  $\log p(w) = \log \sum_z p(w,z)$  increases at each EM iteration.

In this model, the marginal log-likelihood is actually simple to evaluate since you can integrate out each token-level z in turn as you did in the per-sentence log-likelihood question. Please use the marginal probability equation to calculate the corpus log-likelihood and report it at each iteration. This will help verify your algorithm is working correctly. (But MLL calculation may be a bit slow, so you may want to sometimes turn it off during development.)

When reporting marginal log-likelihood, please give the average log-likelihood per token:

$$TokLL = \frac{1}{N_{tok}} \sum_{sentence} \log p(\text{words in sentence})$$

where  $N_{tok}$  is the number of probabilistic word generation events (number of tokens in the corpus plus number of END symbols). Note: perplexity is  $\exp(-TokLL) = \frac{1}{\exp(TokLL)}$ .

**Q4.5 (2 point):** Our dataset has approximately V = 50000. Therefore we should expect per-token average log probability to be higher than -10.8. Why?

**Q4.6** (20 points): Once you feel relatively confident that training is working correctly, run EM for 20 iterations with K=3 latent classes. Run this for three different random initializations, and save the per-iteration log-likelihoods each time. Plot all three curves on the same graph. Discuss the convergence behavior of EM.

Tip: You could save log-likelihood trajectories in global variables or something, or, you could simply print out the likelihoods at each iteration, save the output of your script, then string-munge them back out for the plot.

**Q4.7** Optional extra credit. Prove the statement in Q4.4 that marginal likelihood increases in every EM iteration. You will want to analyze how EM optimizes the variational objective  $J(Q, \theta) = \mathbb{E}_Q[\log p_{\theta}(w, z) - \log Q(z)]$ .

## Q5: Analyzing one model.

20 EM iterations with K=3 should be enough to get the "colorless" example to work. We've noticed that it doesn't always work, so if necessary, you may have to find a model run that does. Once you have this model, conduct some more fine-grained analysis of it. (Jupyter notebook may be a convenient approach for this.)

**Q5.1** (**4 points**): Let's compare to Pereira 2000, page 1245. Calculate and show the following two quantities. (We'll use the American English spelling of "colorless" to match our corpus.)

 $\log p$ (colorless green ideas sleep furiously)  $\log p$ (furiously sleep ideas green colorless)

What is the probability ratio inferred by your model? How does it compare to Pereira's?

**Q5.2** (1 point): Why can this model infer meaningful probabilities while the classic bigram Markov model cannot?

**Q5.3** (**5 points**): Try at least 3 other examples of a well-formed sentence and a nonsensical reordering. Which ones does the model get right or wrong? (Use sentences where all words are in-vocabulary).

**Q5.4** (10 points): Find an example of a well-formed sentence and a reordering of it where you think the model gets the directionality wrong (or use one of the above). Explain what's going on. Develop a falsifiable hypothesis about your model and its linguistic properties, then develop a few more examples to test it on. Criticize and suggest a reformulation or refinement of your hypothesis for future work. How could the model be changed to improve performance on these examples?

### **Q6:** Discussion.

**Q6** (**10 points**): The model you implemented is capable of evaluating sentence probability. Is this the same thing as grammatical well-formedness? Or some other linguistic thing?

For each side of this question, present at least one major argument. We expect a few paragraphs of thoughtful discussion as the answer to this question. Back up your arguments with examples. Beyond the Pereira (2000) reading, for some ideas see:

- Joe Pater's blog post: http://blogs.umass.edu/comphon/2015/06/02/wellformedness-probability/
- Peter Norvig's comments on Noam Chomsky's 2011 remarks: http://norvig.com/chomsky.html

## Implementation tricks and tips:

#### EM:

- First focus on defining the data structures and randomly initializing the parameters.
- Then get EM to run on just 1 iteration on a subset of sentences.
- Then get it to run for 1 iteration on all the sentences.
- Then try more iterations.
- Use just K = 3, since a low K is faster.
- Try a number of random restarts: EM is initialization-dependent so they will give different results.
- To tell whether it's working: at each iteration try printing out
  - The probabilities for a few sample sentences, like the "colorless" examples.
  - The corpus per-token log likelihood, though this is slow.

- Print out top-10 most probable output words per class, excluding globally common words. The problem with outputting the most probable words in this model is that it doesn't learn to put function words in their own class; instead, they sit in all the classes at the top of each distribution. But if you dig a little deeper you can see this better; only rank among words that are outside the set of 1000 most frequent words in the corpus. You should be able to see different syntactic categories emerge during learning.
  - \* Code tip: numpy.argsort(-probs)[:10] is an easy way to get the indexes of the 10 highest entries, in descending rank order, from a vector (1d array) of probabilities.
  - \* Another method is to rank words by the ratio  $\frac{p(w|z=k)}{p(w)}$ , instead of just  $p(w \mid z=k)$ , a.k.a. "exponentiated pointless mutual information". This automatically focuses on words especially unique to the class. Unfortunately, PMI overly aggressively prefers rare words and the words at the top of this ratio-ranked list will be extremely rare ones. If you try this, you'll have to add a filter to only include words with a count of at least 50 or 100 or something.

#### • Other notes:

- Speed: Our naive Python implementation, which iterates over each token position (this is slow since Python is slow!), takes less than a minute per iteration.
- Data structures: There should be two big matrices of parameters: the "transitions" p(class | w<sub>prev</sub>) and the "emissions" p(w<sub>next</sub> | class). You could implement them in two-dimensional numpy arrays each. If so, you will need to track the vocabulary: a two-way map between word-strings and a unique integer index for each wordtype. We usually do this with a Python list (for integer → wordstring) and a Python dict (for wordstring → integer). If you don't like this, you could use Python dicts for everything instead (e.g. use a few word → prob maps), but it will be slower.

#### Other tips:

- Python 2.7 is still often used in NLP/ML. If you're using it, don't forget "from \_\_future\_\_ import division".
- We recommend writing functions in a Python file, then running them from a Jupyter notebook located in the same directory. You can write test code to call specific functions. If the file is called "lm.py", then just add this little line in the top of a cell: import lm;reload(lm)
  - then call the various functions from the module "lm" that you want. Every time you re-execute the cell, you'll get the current code.
- One issue with module reloading is that object methods don't work well. It's a good idea to make a model class with all the data structures for the model. And maybe even to put initialization code as methods on that class. But if you put training or inference code as methods on that class, when you call reload(lm), your already-existing model objects will be stuck with the old methods, not the new ones. This will be confusing. But you don't want to re-train your model every time you change your code; maybe you want to debug your log likelihood function, for example. A better solution is to put important code as standalone functions in the file which simply take a model object as one of their arguments. This is the "dumb objects that are just a pile of data structures" approach to programming, as opposed to "smart objects that have functionality inside of them".

• If you are bored waiting for one iteration to finish, try printing out a progress indicator with e.g.:

```
for si,sentence in enumerate(sentences):

if si % 1000 == 0: sys.stderr.write(".")

[...rest of code...]
```